REMARKS

This Amendment cancels claims 2-4 and amends claim 1, 7 and 13 in accordance with the original disclosure. In view of these amendments and the following remarks, Applicants submit that all of pending claims 1, 5-20 and 23-32 are now in condition for allowance.

Unexamined Claims

As discussed with Examiner Pak in telephone conversations on August 19, 2005 and September 15, 2005, elected claims 23-32 were not examined in the outstanding Office Action. The Examiner instructed Applicants to respond to the rejections of claims 1-20 and stated that claims 23-32 would be examined in a subsequent, non-final, Office Action. Therefore, Applicants base the above amendments and the following remarks on currently examined claims 1-20.

Claim Objections

Claims 7 and 13 stand objected to as being dependent upon a rejected base claim but would be allowable if rewritten in independent form. As set forth above, Applicants have rewritten claims 7 and 13 in independent form.

Rejections Under 35 U.S.C. § 102(b)

Claims 1-6, 8-12 and 14-20 stand rejected under 35 U.S.C. 35 U.S.C. § 102(b) for asserted anticipation by van den Beukel et al. (Applicants believe that the Examiner is referring to the van den Beukel et al. disclosure, hereinafter "van den Beukel"). The Examiner states that van den Beukel discloses chimeric receptors comprising rat alpha and Drosophila alpha subunits, chicken beta 2 subunits; and nucleic acids encoding the above receptors expressed in oocytes. Additionally, the Examiner states that the chimeric DNA were subcloned into pcDNA3 which inherently requires transformation of the host cell.

Applicants submit that the chimeric subunits disclosed by van den Beukel can be distinguished from the present invention with respect to the sequence segments which are replaced with insect sequences. Specifically, attached herewith is a sequence comparison which identifies the regions in van den Beukel's most extended replacement (designated S7.1, page 1032) in italicized type. The region defined by SEQ ID NO: 1 in Torpedo alpha 1 is written in bold and italicized. Thus, it is clear that the modified acetylcholine receptor subunit of the claimed invention comprising an α subunit of a vertebrate acetylcholine receptor in which the entire

region, which is homologous with the amino acid sequence shown in SEQ ID NO: 1, has been replaced by the corresponding region of an α subunit of an insect acetylcholine receptor is not anticipated by van den Beukel.

Applicants also respectfully submit that the Examiner is incorrect that van den Beukel discloses receptors formed from chimeric subunits containing a Beta 2 subunit. Nowhere in van den Beukel is such a description taught or suggested. Rather, van den Beukel discloses a hybrid receptor consisting of Drosophila $\alpha 2$ subunits and chick $\beta 2$ subunits (e.g., page 1032, Fig. 1). It is therefore clear that this receptor does not contain <u>chimeric</u> subunits.

Furthermore, the modified receptor subunits produced by van den Beukel are not capable of a functional response to typical acetylcholine receptor ligands such as acetylcholine or physostigmine (see page 1032, Methods and Results: "Neither ACh, Phytoestrogen nor teramethylammonium was able to induce an ion current in any of the chimeras;" also see page 1032, Conclusions: "Lack of function of the alpha7-SAD chimeras appears to be due to deficient coupling of agonist binding to gating of the ion channel..".). The chimeras disclosed by van den Beukel therefore are not suitable for a test method with which it is possible to find compounds which, as modulators, in particular as agonists or antagonists, alter the conduction properties of insect nicotinic receptors.

In contrast to van den Beukel, the chimeric subunits of the claimed invention assemble with β subunits, forming a receptor that responds with increased ion conductance to the binding of acetylcholine or imidacloprid, not only in Xenopus oocytes, but also in cell lines such as Sf9 (Examples 2 and 3 of the application).

Conclusion

In view of the above amendments and remarks, Applicants respectfully request allowance of all pending claims 1, 5-20 and 23-32.

Respectfully submitted,

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Attachment 2

		* 20 * 40 *	60	
a1_Human 57	:	VAKLFKDYSSVVRPVEDHRQVVEVTVGLQLIQLINVDEVNQIVTTNVRLKQGDMV	/DL	:
a1_Torpedo 51	:	VANLLENYNKVIRPVEHHTHFVDITVGLQLIQLISVDEVNQIVETNVRLRQ		:
a4_Chick 51	:	LKKLFSGYNKWSRPVANISDVVLVRFGLSIAQLIDVDEKNQMMTTNVWVKQ		:
a2_Drosophila 51	:	YDDLLSNYNRLIRPVSNNTDTVLVKLGLRLSQLIDLNLKDQILTTNVWLEH		:
a2_Heliothis	:	YDDLLSNYNRLIRPVDKNNNTVLVKLGLRLSQLIDLNLKDQILTTNVWLEH		:
a3_Heliothis 51	:	YDDLLSNYNRLIRPVTNVSDILTVRLGLKLSQLMEVNLKNQVMTTNLWVEQ		:
a2_Myzus 51	:	YDDLLSNYNRLIRPVGNNSDRLTVKMGLKLSQIIEVNLRNQIMTTNVWVEQ		:
a3_Drosophila	:	YDDLLSNYNKLVRPVVNVTDALTVRIKLKLSQLIDVNLKNQIMTTNLWVEQ		:
al_Heliothis 51	:	YDDLLSNYNKLVRPVLNVSDALTVRIKLKLSQLIDVNLKNQIMTTNLWVEQ		:
a3_Myzus 51	:	YDDLLSNYNKLVRPVLNNTDPLPVRIKLKLSQLIDINLKNQIMTTNLWVEQ		:
a7_Rat 54	:	RRLYKELVKNYNPLERPVANDSQPLTVYFSLSLLQIMDVDEKNQVLTTNIWLQM		:
		* 80 * 100 * 1	20	
al_Human 117	:	* 80 * 100 * 1 PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAD	.20 GD	:
117 al_Torpedo	:		GD	:
117	:	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAD	GD	:
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila	: : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAC)GD)GD)GD	:
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis	: : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNAD	OGD OGD OGD	: : :
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis	: : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNAD	OGD OGD OGD OGE	: : : : :
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus	: : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNADEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAD	OGD OGD OGD OGE OGE	: : : :
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila	: : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNADEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNADEWEDHKFKWDPLEYGGVKELYVPSEHIWLPDIVLYNNAD	OGD OGD OGD OGE OGE OGN	:
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila 92 a1_Heliothis	: : : : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNADEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNADKWFDYKLQWNPDDYGGVEMLYVPSEHIWLPDIVLYNNAD	OGD OGD OGD OGE OGE OGN OGN	:
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila 92 a1_Heliothis 92 a1_Heliothis	: : : : : : :	PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNADEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAD	OGD OGD OGD OGD OGG OGG OGG OGG OGG OGG	: : : : : : : : : : : : : : : : : : : :
117 a1_Torpedo 92 a4_Chick 92 a2_Drosophila 92 a2_Heliothis 92 a3_Heliothis 92 a2_Myzus 92 a3_Drosophila 92 a1_Heliothis		PRPSCVTLGVPLFSHLQNEQWVDYNLKWNPDDYGGVKKIHIPSEKIWRPDLVLYNNADQWIDVRLRWNPADYGGIKKIRLPSDDVWLPDLVLYNNADEWHDYKLRWDPQEYENVTSIRIPSELIWRPDIVLYNNADEWQDHKFKWDPSEYGGVTELYVPSEHIWLPDIVLYNNAD	OGD OGD OGD OGC OGG OGG OGG OGG OGG OGG OGG OGG OGG	: : : : :



		*	140	*	160	*	180	
al_Human 177	:	FAIVKFTKVLLQY		FKSYCEIIVTI		MKLGTWTYD		:
al_Torpedo	:	FAIVHMTKLLLDY	TGKIMWTPPAI)	FKSYCEIIVTI	HFPFDQQNCT	MKLGIWTYD	GTKVSIS	:
a4_Chick 152	:	FAVTHLTKAHLFY	DGRIKWMPPAI:	YKSSCSIDVT	FFFFDQQNCK	MKFGSWTYD	KAKIDLV	:
a2_Drosophila	:	YVVTTMTKAILHY	rgkvvwtppai!	FKSSCEIDVRY	(FPFDQQTCF	MKFGSWTYD	GDQIDLK	:
a2_Heliothis	:	YVVTTMTKAVLHH	rgkvlwtppai:	FKSSCEIDVRY	/FPFDQQTCF	LKFGSWSYD	GDQIDLK	:
a3_Heliothis	:	YEVTLMTKATLKY	rgevnwkppai:	YKSSCEINVEY	FPFDEQTCF	MKFGSWTYN	GAQVDLK	:
a2_Myzus 152	:	YEVTIMTKAILHY!	rgkvvwkppai)	KSFCEINVEY	(FPFDEQTCS	MKFGSWTYD	GYMMDLR	:
a3_Drosophila	:	FEVTLATKATLNY	rgrvewrppai)	YKSSCEIDVEY	(FPFDEQTCV	MKFGSWTYD	GFQVDLR	:
al_Heliothis	:	FEVTLATKATLNY	TGRVEWRPPAI)	KSSCEIDVEY	(FPFDQQTCV	MKFGSWTYD	GFQVDLR	:
a3_Myzus 152	:	FEVTLATKAMLHYS	GRVEWKPPAI	KSSCEIDVER	FFPFDEQTCV	MKFGSWTYD	GFQVDLR	:
a7_Rat 155	:	FDATFHTNVLVNAS	GG <u>HCQYLPPGI</u>	FKSSCYIDVR	VFPFDVQQCK	LKFGSWSYG	<u>GWSLDLQ</u>	:
	-	*	200	*	220	*	240	
a1_Human 227	:	* PESDQP		* GEWVIKESRGW		* PDTPYLDIT		: ,
	:	PESDQP	DLSNFMESO	* GEWVIKESRGW GEWVMKDYRGW	KHSVTYSCC		YHFVMQR	: .
227 al_Torpedo	:	PESDRP	DLSNFMESO	SEWVMKDYRGW	KHSVTYSCC	PDTPYLDIT	YHFVMQR YHFIMQR	-
227 al_Torpedo 202 a4_Chick	:	PESDRP	DLSNFMESO	GEWVMKDYRGN GEWVIINAVGN	KHSVTYSCC KKHWVYYTCC NYNSKKYECC	PDTPYLDIT	YHFVMQR YHFIMQR YSFIIRR	:
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila		PESDRP	DLSNFMESO DLSTFMESO DQLDYWESO	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE	KHSVTYSCC KHWVYYTCC YNSKKYECC CHEKYYPCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR	:
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis	:	PESDRPSMHSHV	DLSNFMESO DLSTFMESO DQLDYWESO IGIDLREYYPSV /GIDLREYYPSV	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE VEWDILGVPAE	KHSVTYSCC KKHWVYYTCC YNSKKYECC CRHEKYYPCC CRHERYYPCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF QEP-YPDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR	:
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis	:	PESDRP SMHSHV HISQKNDKDNKVE	DLSNFMESO DLSTFMESO DQLDYWESO IGIDLREYYPSV /GIDLREYYPSV	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE VEWDILGVPAE VEWDILEVPAT	KHSVTYSCC KHWVYYTCC YNSKKYECC CRHEKYYPCC CRHERYYPCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF QEP-YPDIF PEP-FSDIT	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FNITLRR FKLTMRR	:
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus	:	PESDRP SMHSHV HISQKNDKDNKVE HINQKKGDMVDV	DLSNFMESO DLSTFMESO DQLDYWESO GIDLREYYPSV /GIDLREYYPSV /GIDLSEFYLSV	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE VEWDILGVPAE VEWDILEVPAT	KHSVTYSCC KHWVYYTCC KYNSKKYECC CRHEKYYPCC CRHERYYPCC CRNEEYYPCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF QEP-YPDIF PEP-FSDIT EEP-YLDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FNITLRR FKLTMRR FNITLRR	: : :
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus 210 a3_Drosophila	:	PESDRP SMHSHV HISQKNDKDNKVET HINQKKGDMVDV HMDQSP-GSSLVHV HISQAP-DSDVIEV	DLSNFMESODLSTFMESODQLDYWESO GIDLREYYPSV /GIDLREYYPSV /GIDLSEFYLSV /GIDLQDYYLSV	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE VEWDILGVPAE VEWDILEVPAT VEWDIMGVPAV	KHSVTYSCC KHWVYYTCC KYNSKKYECC CRHEKYYPCC CRHERYYPCC CRNEEYYPCC KRHEKFYVCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF QEP-YPDIF PEP-FSDIT EEP-YLDIF	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR FNITLRR FNITLRR	: : : : : : : : : : : : : : : : : : : :
227 al_Torpedo 202 a4_Chick 201 a2_Drosophila 211 a2_Heliothis 209 a3_Heliothis 210 a2_Myzus 210 a3_Drosophila 210 a1_Heliothis	:	PESDRP SMHSHV HISQKNDKDNKVET HINQKKGDMVDV HMDQSP-GSSLVHV HISQAP-DSDVIEV	DLSNFMESODLSTFMESODQLDYWESO GIDLREYYPSV /GIDLREYYPSV /GIDLSEFYLSV /GIDLQDYYLSV /GVDLSEFYTSV	GEWVMKDYRGW GEWVIINAVGN VEWDILGVPAE VEWDILEVPAT VEWDIMGVPAV VEWDILEVPAV	KHSVTYSCC WKHWVYYTCC WYNSKKYECC CRHEKYYPCC CRHERYYPCC WRNEEYYPCC WRNEKFYVCC WRNEKFYTCC	PDTPYLDIT TEI-YPDIT AEP-YPDIF QEP-YPDIF PEP-FSDIT EEP-YLDIF DEP-YLDIT	YHFVMQR YHFIMQR YSFIIRR FNITLRR FNITLRR FKLTMRR FNITLRR FNITLRR FNITLRR	: : : : : : : : : : : : : : : : : : : :